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<!--StartFragment-->RESULT 1
US-09-828-217-1
; Sequence 1, Application US/09828217
; Patent No. US20010051710A1
;   GENERAL INFORMATION:
;       APPLICANT: HIRANO, TOSHIO
;               KAISHO, TSUNEYASU
;   TITLE OF INVENTION: MEMBRANE PROTEIN POLYPEPTIDE HAVING
;                       PRE-B CELL GROWTH-SUPPORTING ABILITY AND A GENE THEREOF
;   NUMBER OF SEQUENCES: 2
;   CORRESPONDENCE ADDRESS:
;       ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
;               P.C.
;       STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400
;       CITY: ARLINGTON
;       STATE: VA
;       COUNTRY: USA
;       ZIP: 22202
;   COMPUTER READABLE FORM:
;       MEDIUM TYPE: Floppy disk
;       COMPUTER: IBM PC compatible
;       OPERATING SYSTEM: PC-DOS/MS-DOS
;       SOFTWARE: PatentIn Release #1.0, Version #1.30
;   CURRENT APPLICATION DATA:
;       APPLICATION NUMBER: US/09/828,217
;       FILING DATE: 09-Apr-2001
;       CLASSIFICATION: <Unknown>
;   PRIOR APPLICATION DATA:
;       APPLICATION NUMBER: 09/182,563
;       FILING DATE: <Unknown>
;       APPLICATION NUMBER: PCT/JP94/01732
;       FILING DATE: 14-OCT-1994
;       APPLICATION NUMBER: JP 5-281622
;       FILING DATE: 15-OCT-1993
;   ATTORNEY/AGENT INFORMATION:
;       NAME: OBLON, NORMAN F.
;       REGISTRATION NUMBER: 24,618
;       REFERENCE/DOCKET NUMBER: 7625-001-0 PCT
;   TELECOMMUNICATION INFORMATION:
;       TELEPHONE: 703-413-3000
;       TELEFAX: 703-413-2220
;   INFORMATION FOR SEQ ID NO: 1:
;       SEQUENCE CHARACTERISTICS:
;           LENGTH: 180 amino acids
;           TYPE: amino acid
;           STRANDEDNESS: single
;           TOPOLOGY: linear
;       MOLECULE TYPE: peptide
;       SEQUENCE DESCRIPTION: SEQ ID NO: 1:
US-09-828-217-1

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Query Match          100.0%;  Score 889;  DB 3;  Length 180;
Best Local Similarity 100.0%;  Pred. No. 2.9e-77;
Matches 180;  Conservative 0;  Mismatches 0;  Indels 0;  Gaps 0;

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Qy      1 MASTSYDYCRVPMEDGDKRCKLLLIGIGILVLLIIVILGVPLIIFTIKANSEACRDGLRAV 60
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Db      1 MASTSYDYCRVPMEDGDKRCKLLLIGIGILVLLIIVILGVPLIIFTIKANSEACRDGLRAV 60

Qy      61 MECRNVTHTLLQQLTEAQKGFQDVEAQAATCNHTVMALMASLDAEKAQGQKKVEELEGEI 120
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Db 61 MECRNVTHLLQQELTEAQKGFQDVEAQAATCNHTVMALMASLDAEKAQGQKKVEELEGEI 120

Qy 121 TTLNHKLQDASAEVERLRRENQVLSVRIADKKYYPSSQDSSSAAAPQLLIVLLGLSALLQ 180

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Db 121 TTLNHKLQDASAEVERLRRENQVLSVRIADKKYYPSSQDSSSAAAPQLLIVLLGLSALLQ 180

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